Supplementary Information:

Table S1. The Conserved Motifs/Residues of RAF Family Kinases Regulating Their Activity

Raf Isoform	NtA motif	Catalytic spine mutation	Regulatory spine mutation	Central RH alteration in dimer interface	APE motif
ARAF	299 SGYY 302	V324F	L358M	R362H	474AAE
BRAF	446SSDD449	V471F	L505M	R509H	621APE
CRAF	338SSYY 341	V363F	L397M	R401H	₅₁₃ APE ₅₁₅

Table S1. The Conserved Motifs/Residues of RAF Family Kinases Regulating Their Activity. The key motifs/residues of RAF family kinases involved in this study were listed in this table.

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Table S2. Protein Kinase Mutants With In-frame Deletions of β3-αC Loop in Cancer. To explore all protein kinase mutants with variable β 3-αC loop deletions in cancer genomes, we we interrogated the ICGC (International Cancer Genome Consortium) database, the cBioportal for Cancer Genomics database, and the COSMIC (Catalogue of Somatic Mutations in Cancer) database. All known mutations including those reported in literatures ¹⁻⁴ were summarized in this table.

Figure S1

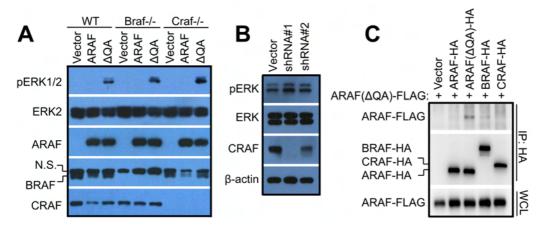


Figure S1 (related to Figure 1). ARAF mutant with Q347A348 deletion is activated by elevated homodimerization but not heterodimeization.

A, ARAF(ΔQA) and its wild-type counterpart were stably expressed in RAF-knockout fibroblasts, and their activity was measured by anti-phosphoERK1/2 immunoblot. ARAF(ΔQA) activates MEK-ERK signaling in wild-type, BRAF-/-, and CRAF-/- MEFs. B, the CRAF knockdown does not affect the activity of ARAF(ΔQA) in BRAF-/- WEFs. The activity of ARAF(ΔQA) in BRAF-/- with or without shRNAs against CRAF was measured as phosphoERK by immunoblot. C, ARAF(ΔQA) hardly heterodimerizes with wild-type ARAF, BRAF and CRAF. The wild-type RAF molecules were coexpressed with ARAF(ΔQA) in 293T cells, and the coimmunoprecipitation assay was carried out as described before⁵. All images are representative of at least three independent experiments.

Figure S2

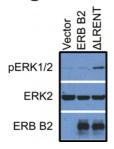


Figure S2. An ERB B2 mutant with β 3- α C loop deletion has constitutive activity.

The ERB B2 mutant with in-frame $\beta3-\alpha C$ loop deletion was expressed in 293T cells and its activity was measured by anti-phosphoERK1/2 immunoblot. All images are representative of at least three independent experiments.

Figure S3

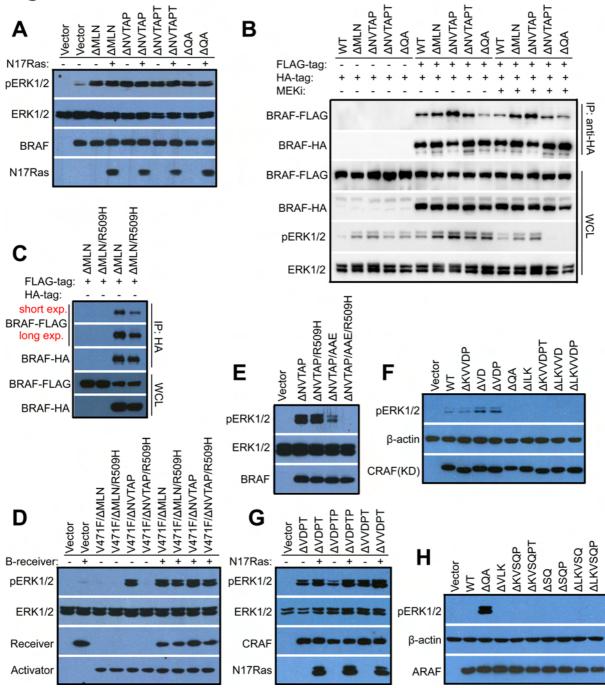


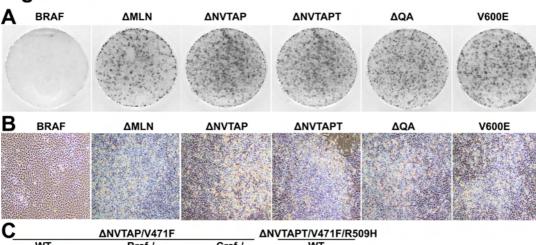
Figure S3 (related to Figure 4). RAF kinase is activated by in-frame β 3- α C loop deletions through enhanced homodimerization.

- (A) BRAF mutants with in-frame $\beta3-\alpha C$ loop deletions exhibit a RAS-independent activity. The BRAF mutants were expressed with or without N17RAS in 293T cells and their activity was measured by anti-phosphoERK1/2 immunoblot.
- (B-C) BRAF mutants with in-frame $\beta 3-\alpha C$ loop deletions have increased albeit variable dimer affinity. The dimer affinity of BRAF mutants in 293T transfectants treated with or without MEK inhibitor was measured by co-immunoprecipitation as in Figure 1E.
- (D) The central R509H alteration in dimer interface is not able to abolish the allosteric activity of Catalytic spine-fused BRAF mutants with a high dimer affinity toward BRAF-receiver. The RAF co-activation assay was carried out as described before 5.6.

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- (E) The combined alteration of APE motif and the central Arg in dimer interface abolishes the activity 111 of BRAF mutant with a high dimer affinity. The BRAF mutants were expressed in 293T cells and their activity was measured by anti-phosphoERK1/2 immunoblot.
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Figure S4



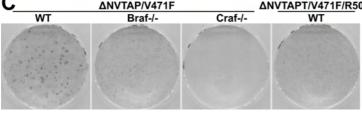
(F-G) In-frame β3-αC loop deletions turn on CRAF independent of RAS. The activity of CRAF mutants

(H) ARAF is activated by ΔQA but not other deletions in $\beta 3-\alpha C$ loop. The activity of ARAF mutants

expressed in 293T cell with or without N17RAS was measured as in (A).

All images are representative of at least three independent experiments.

expressed in 293T cells was measured as in (A).



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Figure S4 (related to Figure 5). BRAF mutants with in-frame β3-αC loop deletions have a strong 123 oncogenic potential. 124

(A-B) BRAF mutants with in-frame β3-αC loop deletions transform both BRAF-knockout and CRAFknockout immortalized fibroblasts. A, foci formation assay of immortalized braf /- MEFs that express BRAF mutants. B, cellular morphology images of immortalized craf^{-/-} MEFs that express BRAF (C) The transforming ability of kinase-dead BRAF(ΔNVTAP/V471F) depends on endogenous RAF

molecules and dimerization. Foci formation assay of immortalized MEFs that express BRAF mutant

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was carried out as in Figure 1C. All images are representative of at least three independent experiments.

Figure S5

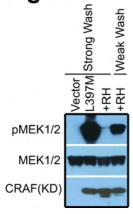


Figure S5 (related to Figure 6). The constitutively active CRAF R-spine mutant with a low dimer affinity retains partial catalytic activity in vitro upon purification with a gentle wash of PBS. CRAF mutants were expressed in 293T cells and purified by immunoprecipitation with either a strong wash of RIPA buffer (1% NP-40) or a gentle wash of PBS. Then their activity was measured by in vitro kinase assay as in Figure 6.

All images are representative of at least three independent experiments.



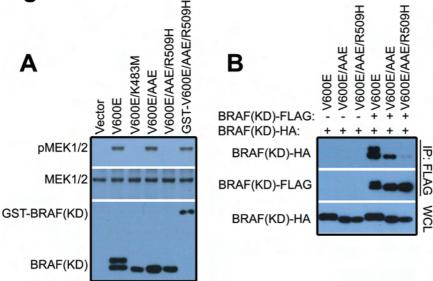


Figure S6 (related to Figure 7). The BRAF(V600E) mutant with non-canonical APE motif has constitutive activity but is sensitive to RH alteration in dimer interface.

(A) The BRAF(V600E) mutant with non-canonical APE motif can phosphorylate MEK as well as its prototype, which is blocked by further RH alteration in dimer interface. BRAF mutants were expressed in 293T cells and the phospho-MEK1/2 was detected by immunoblot. The kinase-dead BRAF(V600E) mutant, BRAF(V600E/K483M) was used as control.

(B) The non-canonical APE motif decreases the dimer affinity of BRAF(V600E) but can not completely abolish its dimerization. FLAG- and HA-tagged BRAF mutants were co-expressed in 293T cells that were lyzed with RIPA buffer containing 0.1% NP-40, and the co-immunoprecipitation assay was carried out with a gentle wash of PBS as in Figure S5.

All images are representative of at least three independent experiments.

Supplementary Materials and Methods

To knockdown CRAF in braf MEFs that stably express ARAF(ΔQA), shRNAs were designed by using a website software (http://katahdin.cshl.org/siRNA/RNAi.cgi?type=shRNA), and the lentiviral vectors were constructed by using traditional molecular cloning methods. The targeting sequences were listed as follow:

shRNA#1 5- GGAATGGAATGAGCTTACA-3 shRNA#2 5- GGAATGAGCTTACATGACT-3

Reference

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